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NOTICE OF ALLOWANCE AND FEE(S) DUE

65565	

08/11/2008

SUGHRUE-265550 2100 PENNSYLVANIA AVE. NW WASHINGTON, DC 20037-3213 EXAMINER

LEE, JAE W

ART UNIT PAPER NUMBER

1656

DATE MAILED: 08/11/2008

APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
10/537,767	06/06/2005	Yuki Endo	Q88255	5081

TITLE OF INVENTION: PROTEIN WHICH BINDS TO AKT2

APPLN. TYPE	SMALL ENTITY	ISSUE FEE DUE	PUBLICATION FEE DUE	PREV. PAID ISSUE FEE	TOTAL FEE(S) DUE	DATE DUE
nonprovisional	NO	\$1440	\$300	\$0	\$1740	11/12/2008

THE APPLICATION IDENTIFIED ABOVE HAS BEEN EXAMINED AND IS ALLOWED FOR ISSUANCE AS A PATENT. PROSECUTION ON THE MERITS IS CLOSED. THIS NOTICE OF ALLOWANCE IS NOT A GRANT OF PATENT RIGHTS. THIS APPLICATION IS SUBJECT TO WITHDRAWAL FROM ISSUE AT THE INITIATIVE OF THE OFFICE OR UPON PETITION BY THE APPLICANT. SEE 37 CFR 1.313 AND MPEP 1308.

THE ISSUE FEE AND PUBLICATION FEE (IF REQUIRED) MUST BE PAID WITHIN THREE MONTHS FROM THE MAILING DATE OF THIS NOTICE OR THIS APPLICATION SHALL BE REGARDED AS ABANDONED. THIS STATUTORY PERIOD CANNOT BE EXTENDED. SEE 35 U.S.C. 151. THE ISSUE FEE DUE INDICATED ABOVE DOES NOT REFLECT A CREDIT FOR ANY PREVIOUSLY PAID ISSUE FEE IN THIS APPLICATION. IF AN ISSUE FEE HAS PREVIOUSLY BEEN PAID IN THIS APPLICATION (AS SHOWN ABOVE), THE RETURN OF PART B OF THIS FORM WILL BE CONSIDERED A REQUEST TO REAPPLY THE PREVIOUSLY PAID ISSUE FEE TOWARD THE ISSUE FEE NOW DUE.

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B. If applicant claimed SMALL ENTITY status before, or is now claiming SMALL ENTITY status, check box 5a on Part B - Fee(s) Transmittal and pay the PUBLICATION FEE (if required) and 1/2 the ISSUE FEE shown above.

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IMPORTANT REMINDER: Utility patents issuing on applications filed on or after Dec. 12, 1980 may require payment of maintenance fees. It is patentee's responsibility to ensure timely payment of maintenance fees when due.

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Complete and send this form, together with applicable fee(s), to: Mail Mail Stop ISSUE FEE

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CURRENT CORRESPONDENCE ADDRESS (Note: Use Block 1 for any change of address)				s) Transmittal. This rs. Each additional	certifi paper,	icate cannot be used fo	domestic mailings of the or any other accompanying it or formal drawing, must
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							(Depositor's name)
							(Signature)
							(Date)
APPLICATION NO.	FILING DATE		FIRST NAMED INVENTOR		ATTO	RNEY DOCKET NO.	CONFIRMATION NO.
10/537,767 FITLE OF INVENTION	06/06/2005 : PROTEIN WHICH BII	NDS TO AKT2	Yuki Endo			Q88255	5081
APPLN. TYPE	SMALL ENTITY	ISSUE FEE DUE	PUBLICATION FEE DUE	PREV. PAID ISSUE	FEE	TOTAL FEE(S) DUE	DATE DUE
nonprovisional	NO	\$1440	\$300	\$0		\$1740	11/12/2008
EXAM	INER	ART UNIT	CLASS-SUBCLASS				
LEE, J.	AE W	1656	536-023100				
"Fee Address" indi PTO/SB/47; Rev 03-0 Number is required. 3. ASSIGNEE NAME A	ess an assignee is identi h in 37 CFR 3.11. Comp	' Indication form ed. Use of a Customer A TO BE PRINTED ON T	or agents OR, alternative (2) the name of a single registered attorney or a 2 registered patent attoo listed, no name will be THE PATENT (print or type data will appear on the path a substitute for filing and (B) RESIDENCE: (CITY)	e firm (having as a regent) and the name neys or agents. If norinted. e) tent. If an assigners is a signer signment.	s of up o nam	entified below, the do	cument has been filed for
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	are submitted: fo small entity discount p # of Copies	permitted)	 Payment of Fee(s): (Plea A check is enclosed. Payment by credit care The Director is hereby overpayment, to Depos 	d. Form PTO-2038 authorized to charg	is atta	ched. required fee(s), any def	
a. Applicant claims	tus (from status indicated s SMALL ENTITY statu	is. See 37 CFR 1.27.	☐ b. Applicant is no long				
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This collection of inform an application. Confident submitting the completed his form and/or suggesti Box 1450. Alexandria, V	ation is required by 37 C tiality is governed by 35 I application form to the ons for reducing this but irginia 22313-1450 DC	FR 1.311. The information U.S.C. 122 and 37 CFR USPTO. Time will vary to be sent to the DNOT SEND FEES OR 6	on is required to obtain or re 1.14. This collection is esti- depending upon the indive e Chief Information Office COMPLETED FORMS TO	etain a benefit by th mated to take 12 m idual case. Any con r, U.S. Patent and T D THIS ADDRESS	e publ inutes nment raden SENT	ic which is to file (and to complete, including s on the amount of tin lark Office, U.S. Depa of TO: Commissioner f	by the USPTO to process) g gathering, preparing, and he you require to complete rtment of Commerce, P.O. for Patents, P.O. Box 1450.

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10/537,767	06/06/2005	Yuki Endo	Q88255 5081	
65565 7	590 08/11/2008		EXAM	IINER
SUGHRUE-265	550	LEE, J	AE W	
2100 PENNSYLVANIA AVE. NW			ART UNIT	PAPER NUMBER
WASHINGTON,	WASHINGTON, DC 20037-3213			
		DATE MAILED: 08/11/200	8	

Determination of Patent Term Adjustment under 35 U.S.C. 154 (b)

(application filed on or after May 29, 2000)

The Patent Term Adjustment to date is 143 day(s). If the issue fee is paid on the date that is three months after the mailing date of this notice and the patent issues on the Tuesday before the date that is 28 weeks (six and a half months) after the mailing date of this notice, the Patent Term Adjustment will be 143 day(s).

If a Continued Prosecution Application (CPA) was filed in the above-identified application, the filing date that determines Patent Term Adjustment is the filing date of the most recent CPA.

Applicant will be able to obtain more detailed information by accessing the Patent Application Information Retrieval (PAIR) WEB site (http://pair.uspto.gov).

Any questions regarding the Patent Term Extension or Adjustment determination should be directed to the Office of Patent Legal Administration at (571)-272-7702. Questions relating to issue and publication fee payments should be directed to the Customer Service Center of the Office of Patent Publication at 1-(888)-786-0101 or (571)-272-4200.

	Application No.	Applicant(s)
	10/537,767	ENDO ET AL.
Notice of Allowability	Examiner	Art Unit
	JAE W. LEE	1656
The MAILING DATE of this communication apperall claims being allowable, PROSECUTION ON THE MERITS IS herewith (or previously mailed), a Notice of Allowance (PTOL-85) NOTICE OF ALLOWABILITY IS NOT A GRANT OF PATENT RI	ears on the cover sheet w (OR REMAINS) CLOSED or other appropriate comm GHTS. This application is	ith the correspondence address in this application. If not included nunication will be mailed in due course. THIS
1. This communication is responsive to <u>06/20/2008</u> .		
2. ☑ The allowed claim(s) is/are <u>1-5</u> .		
 3. Acknowledgment is made of a claim for foreign priority ur a) All b) Some* c) None of the: 1. Certified copies of the priority documents have 2. Certified copies of the priority documents have 3. Copies of the certified copies of the priority documents 	been received. been received in Applicati	on No
International Bureau (PCT Rule 17.2(a)).	cuments have been receive	ed in this national stage application from the
* Certified copies not received:		
Applicant has THREE MONTHS FROM THE "MAILING DATE" noted below. Failure to timely comply will result in ABANDONN THIS THREE-MONTH PERIOD IS NOT EXTENDABLE. 4. A SUBSTITUTE OATH OR DECLARATION must be subm INFORMAL PATENT APPLICATION (PTO-152) which give	IENT of this application. itted. Note the attached EX	AMINER'S AMENDMENT or NOTICE OF
5. CORRECTED DRAWINGS (as "replacement sheets") mus	. ,	
(a) ☐ including changes required by the Notice of Draftspers		w (PTO-948) attached
1) ☐ hereto or 2) ☐ to Paper No./Mail Date	-	(
(b) ☐ including changes required by the attached Examiner's Paper No./Mail Date Identifying indicia such as the application number (see 37 CFR 1	s Amendment / Comment o	the drawings in the front (not the back) of
each sheet. Replacement sheet(s) should be labeled as such in the first of the sheet. Before and/or INFORMATION about the deposit attached Examiner's comment regarding REQUIREMENT	sit of BIOLOGICAL MAT	ERIAL must be submitted. Note the
Attachment(s)	5 🗖 Notice of I	oformal Patant Application
 Notice of References Cited (PTO-892) D Notice of Draftperson's Patent Drawing Review (PTO-948) 	<u> </u>	nformal Patent Application Summary (PTO-413),
3. ☐ Information Disclosure Statements (PTO/SB/08),	Paper No	./Mail Date s Amendment/Comment
Paper No./Mail Date 4. Examiner's Comment Regarding Requirement for Deposit		s Statement of Reasons for Allowance
of Biological Material	 9.	

EXAMINER'S AMENDMENT

An extension of time under 37 CFR 1.136(a) is required in order to make an examiner's amendment which places this application in condition for allowance. During a telephone conversation conducted on 07/31/2008, Susan Mack requested an extension of time for 2 MONTH(S) and authorized the Director to charge Deposit Account No. 19-4880 the required fee of \$ 450 for this extension and authorized the following examiner's amendment. Should the changes and/or additions be unacceptable to applicant, an amendment may be filed as provided by 37 CFR 1.312. To ensure consideration of such an amendment, it MUST be submitted no later than the payment of the issue fee.

An examiner's amendment to the record appears below. Should the changes and/or additions be unacceptable to applicant, an amendment may be filed as provided by 37 CFR 1.312. To ensure consideration of such an amendment, it MUST be submitted no later than the payment of the issue fee.

Authorization for this Examiner's amendment was given in a telephone interview with Susan Mack on 07/24/2008.

The claims have been amended as follows:

Claim 5, replace "A cell" with "An isolated cell".

The following is an examiner's statement of reasons for allowance:

The prior art of record does not teach or fairly suggest an isolated polynucleotide encoding a polypeptide which comprises the amino acid sequence of SEQ ID NO: 2 or

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SEQ ID NO: 4, which binds to Akt-2, the polypeptide sequences of SEQ ID NO: 2 and 4, an expression vector comprising said polynucleotide sequence, and an isolated cell transformed with said expression vector. The utility of the claimed invention resides on the fact that Applicants have discovered that an increased expression of the polypeptides as set forth in SEQ ID NOs: 2 and 4, interferes with the insulin signaling cascade downstream of Akt-2 also known as Protein Kinase B beta (PKBβ) for the activation of GLUT4 (glucose transporter in muscle and adipose tissues) translocation from cytosol to cell membrane for increased sugar uptake. Thus, the claimed polypeptides, polynucleotides and cells have a potential use in the treatment of type 2 diabetes mellitus. The Examiner notes that closest prior art references are: [1] the polynucleotide as set forth in GenBank accession number BC049110 (publicly available on June of 2003), which encodes a protein that comprises SEQ ID NO: 2 except that it contains (a) substitutions at positions 37 and 457, and (b) a deletion from position 120 to position 125 (see the sequence alignment 1 below); [2] the polynucleotide as set forth in Genbank accession number AK128728 (publicly available on September of 2003), which encodes a protein that comprises SEQ ID NO: 4 except that it contains (a) a substitution at position 174 and (b) a deletion from position 1 to position 16 (see the sequence alignment 2 below); and [3] the polynucleotide as set forth in Genbank accession number AX714043 (publicly available on March of 2003), which encodes a protein that comprises all of SEQ ID NO: 4 except for a deletion from position 176 to position 243(splice variant of the polypeptide of SEQ ID NO: 4) (see the sequence alignment 3 below).

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Qу

1. Sequence alignment for BC049110:

US-10-537-767-2 (1-573) x BC049110 (1-2738) $1\ {\tt MetAlaAlaValProProLeuArgAspArgLeuSerPheLeuHisArgLeuProIleLeu}\ 20$ QУ 13 ATGGCAGCTGTGCCGCCCCTGCGGGACCGCTTGAGCTTCTTGCATAGGCTCCCCATCCTG 72 Db 21 LeuLysGlyThrSerAspAspSerIleProCysProGlyTyrLeuPheGluGluIleAla 40 Qy TTGAAGGGGACCTCAGATGATAGCATCCCATGTCCAGGCTACCTGTTTTAAGAGATCGCC 132 Qу 41 LysIleSerHisGluSerLeuGlySerSerGlnCysLeuLeuGluTyrLeuLeuAsnArg 60 Db 133 AAGATTTCCCACGAGTCACTAGGCAGCCAGTGCCTGCTGGAGTACCTCCTGAACCGT 192 LeuAspSerSerGlyHisValLysLeuLysValLeuLysIleLeuLeuTyrLeuCys 80 193 CTGGACAGCAGCTCCGGCCACGTGAAGCTCAAGGTGCTAAAGATCTTGCTTTACCTGTGT 252 Dh 81 GlyHisGlySerSerSerPheLeuLeuIleLeuArgArgAsnSerAlaLeuIleGlnGlu 100 Qy Db 253 GGTCATGGCTCTTCCTCCTCCTCATCCTCAGGAGAAACTCTGCTCTCATCCAAGAA 312 101 AlaThrAlaPheSerGlyProProAspProLeuHisGlyAsnSerLeuTyrGlnLysVal 120 0.7 313 GCCACGGCTTTCTCAGGGCCTCCAGATCCTCTTCACGGAAATAGCTTGTACCAGAAG--- 369 Qу 121 ArgAlaAlaGlnAspLeuGlySerThrLeuPheSerAspAlaValProGlnProPro 140 370 -----GACCTGGGTAGCACCCTGTTCTCAGATGCCGTGCCACAGCCTCCA 414 Db 141 SerGlnProProGlnIleProProProAlaGlyMetGlyAlaGlnAlaArgProLeuSer 160 Qу Db 415 TCGCAGCCACCTCAGATCCCGCCTCCCGCAGGCATGGGCGCCCAGGCCAGACCTCTTAGT 474 161 AlaLeuGlnGlyPheGlyTyrThrLysGluSerSerArgThrGlySerAlaGlyGluThr 180 Qу 475 GCCCTGCAAGGCTTCGGCTACACGAAGGAGCAGCCGGACAGGCTCCGCAGGTGAAACC 534 Db 181 PheLeuSerThrIleGlnArgAlaAlaGluValValAlaAsnAlaValArgProGlyPro 200 Qу Db TTCCTCTCCACCATCCAGAGGGCCGCAGAGGTAGTGGCTAATGCTGTGCGTCCTGGACCT 594 AspAsnProCysThrLysGlyProLeuProTyrGlyAspSerTyrGlnProAlaValThr 220 Qy Db 595 GATAATCCTTGTACCAAGGGACCCTTGCCGTATGGTGATTCCTACCAGCCTGCAGTGACA 654 Q.y 221 ProSerAlaSerHisThrHisProAsnProGlyAsnLeuLeuProGlyAlaIleLeuGly 240 655 CCTTCAGCTAGCCACACACACCCTGGGAATCTACTCCCTGGGGCCATCCTGGGG 714 Db 241 AlaArgAlaValArgHisGlnProGlyGlnAlaGlyGlyGlyTrpAspGluLeuAspSer 260 Qу Db 715 GCCAGAGCTGTGAGACACCAGCCCGGGCAGGCTGGGGCGGCTGGGATGAGCTGGACAGC 774 261 SerProSerSerGlnAsnSerSerCysThrSerAsnLeuSerArgAlaSerAspSerGly 280 Qy 775 AGTCCTAGTTCCCAGAATTCCTCCTGCACCAGCAACCTGAGCAGGGCCTCGGACTCGGGC 834 281 SerArgSerGlySerAspSerHisSerGlyThrSerArgGluProGlyAspLeuAlaGlu 300 835 AGTCGGTCTGGCAGTGACAGCCACTCTGGCACCAGCCGGGAGCCAGGCGACCTGGCAGAA 894 Db

 $\tt 301\ ArgAlaGluAlaThrProProAsnAspCysGlnGlnGluLeuAsnLeuValArgThrVal\ 320$

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Db	895	${\tt AGGGCTGAGGCCACGCCCCCAAATGACTGCCAGCAAGAACTGAATCTAGTGAGGACTGTG}$	954
Qy	321	ThrGlnGlyProArgValPheLeuSerArgGluGluThrGlnHisPheIleLysGluCys	340
Db	955	ACACAGGGGCCACGTGTCTTCCTGAGCCGTGAGGAGCACGCAGCACTTCATCAAAGAGT	1014
Qу	341	GlyLeuLeuAsnCysGluAlaValLeuGluLeuLeuLeuArgGlnLeuValGlyThrSer	360
Db	1015	GGCCTGCTCAACTGTGAGGCAGTGCTGGAGCTGCTCCTGCGCCAGCTGGTCGGGACCAGT	1074
Qу	361	GluCysGluGlnMetArgAlaLeuCysAlaIleAlaSerPheGlySerAlaAspLeuLeu	380
Db	1075	${\tt GAGTGCGAGCAGATGAGGGCGCTGTGTGCCATCGCGTCCTTTGGGAGTGCTGACCTCCTG}$	1134
Qy	381	ProGlnGluHisValLeuLeuLeuCysArgGlnGlnLeuGlnGluLeuGlyAlaGlySer	400
Db	1135	CCTCAGGAGCACGTCCTCCTGTGCCGACAGCAGCTGCAGGAACTTGGCGCGGGCAGC	1194
QУ	401	ProGlyProValThrAsnLysAlaThrLysIleLeuArgHisPheGluAlaSerCysGly	420
Db	1195	CCTGGACCTGTGACCAACAAGCCACCAAGATCCTGAGACATTTTGAAGCCTCCTGTGGA	1254
QУ	421	GlnGlnLeuProThrLeuArgLeuCysAlaGlnProAsnSerAlaAlaAlaProValGly	440
Db	1255	${\tt CAACAGCTCCCTACCCTAAGGCTCTGTGCCCAGCCCAACTCTGCAGCTGCCCCTGTGGGC}$	1314
Qу	441	ProAlaAspLeuLeuThrSerProValProAlaProGlySerGlnValCysLeuGlnPro	460
Db	1315	$\tt CCAGCTGACCTGCCAGCCCGTGCCTGCCCTGGGAGCCAGGTCTTCCTCCAGCCT$	1374
Qу	461	LeuSerSerAlaThrValValProArgSerProValLeuPheProSerProAsnThrLeu	480
Db	1375	$\tt CTCAGCTCCGCCACAGTGGTACCCAGGAGTCCTGTGCTCTTTCCATCCCCCAATACCTTA$	1434
QУ	481	ProProSerAlaLeuGluProSerGluValArgThrGlnLeuValCysSerSerGlu	500
Db	1435	$\verb CCTCCGTCTGCTCTGGAGGAGCCCAGCGAGGTCCGAACCCAATTGGTGTGTTCTAGTGAA $	1494
QY	501	GlnGlyThrGluSerGluGlnArgLeuGluAsnThrAspThrProGluAspSerSerSer	520
Db	1495	CAGGGGACAGAATCTGAGCAGAGGCTGGAGAACACACACA	1554
QУ	521	ProLeuProTrpSerProAsnSerLeuPheAlaGlyMetGluLeuValAlaCysProArg	540
Db	1555	CCGCTCCCGTGGAGTCCCAACTCTTTGTTTGCTGGCATGGAGCTGGTGGCTTGCCCCCGC	1614
Qy	541	LeuProCysHisSerSerGlnAspLeuGlnThrAspLeuGlnLysValThrThrGluAla	560
Db	1615	CTGCCTTGCCACAGCTCGCAGGACCTCCAGACAGATTTACAGAAGGTGACCACAGAAGCT	1674
Qy	561	ProValSerGluProSerAlaPheAlaPheLeuAsnMet 573	
Db	1675	CCGGTTTCAGAGCCATCAGCTTTTGCATTTTTAAACATG 1713	

2. Sequence alignment for AK128728:

US-10-537-767-4 (1-593) x AK128728 (1-3802)

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QУ	57	LeuLeuSerArgLeuHisSerSerSerGlyHisGlyLysLeuLysValLeuLysIleLeu	76
Db	1540	$\tt CTCCTGAGCCGCCTGCACAGCAGCTCCGGCCACGGGAAGCTCAAGGTGCTGAAGATCCTG$	159
QУ	77	LeuTyrLeuCysSerHisGlySerSerPhePheLeuLeuIleLeuLysArgAsnSerAla	96
Db	1600	$\tt CTCTATCTGTGCAGCCACGGCTCCTTCTTCCTGCTCATCCTCAAACGCAACTCTGCC$	165
Qу	97	PheIleGlnGluAlaAlaAlaPheAlaGlyProProAspProLeuHisGlyAsnSerLeu	116
Db	1660	$\tt TTCATCCAGGAAGCTGCAGCTTTTGCAGGGCCCCCAGATCCTCTGCACGGGAACAGCTTG$	171
QУ	117	TyrGlnLysValArgAlaAlaAlaGlnAspLeuGlySerThrLeuPheSerAspThrVal	136
Db	1720	${\tt TACCAGAAGGTTCGCGCGGCGCGCAGGACTTGGGGAGCACCCTGTTCTCGGACACCGTG}$	177
QУ	137	LeuProLeuAlaProSerGlnProLeuGlyThrProProAlaThrGlyMetGlySerGln	156
Db	1780	$\tt TTGCCGCTGGCTCCCCAGCCTCTGGGGACCCCGCCTGCCACAGGCATGGGCTCCCAG$	183
QУ	157	AlaArgProHisSerThrLeuGlnGlyPheGlyTyrSerLysGluHisGlyArgThrGly	176
Db	1840	$\tt GCCAGGCCGCACAGCACCCTCCAGGGTTTCGGCTACAGCAAGGAACACGGCCACACGGGC$	189
QУ	177	SerAlaGlyGluAlaPheLeuSerThrIleGlnLysAlaAlaGluValValAlaSerAla	196
Db	1900	$\tt TCGGCAGGCGAAGCCTTCCTCTCCACCATCCAGAAGGCCGCAGAGGTGGTGGCCAGCGCC$	195
QУ	197	MetArgProGlyProGluSerProSerThrArgArgLeuLeuProArgGlyAspThrTyr	216
Db	1960	ATGCGCCCCGGGCCCGAGAGTCCCAGTACCCGGAGGCTCCTGCCGCGGGGTGACACCTAC	201
QУ	217	GlnProAlaMetMetProSerAlaSerHisGlyProProThrLeuGlyAsnLeuLeuPro	236
Db	2020	CAGCCTGCCATGATGCCTTCAGCCAGCCACGGTCCCCCAACCCTGGGGAACCTACTCCCC	207
QУ	237	GlyAlaIleProGlyProArgAlaValArgHisGlnProGlyGlnAlaGlyGlyTrp	256
Db	2080	$\tt GGGGCCATTCCAGGTCCCCGAGCTGTGAGGCATCAGCCTGGGCAGGGCGGAGGGGGGTGG$	213
QУ	257	AspGluLeuAspSerGlyProSerSerGlnAsnSerSerGlnAsnSerAspLeuSerArg	276
Db	2140	${\tt GATGAGCTGGACAGCGGCCCCAGCTCTCAGAATTCCTCCCAGAACAGCGACCTGAGCAGG}$	219
QУ	277	ValSerAspSerGlySerHisSerGlySerAspSerHisSerGlyAlaSerArgGluPro	296
Db	2200	$\tt GTCTCGGACTCGGCAGTCATTCCGGCAGCGACCAGCCATTCAGGGGCCAGCCGGGAGCCGGAGCCGGGAGCGGAGCGGAGCGGAGCGGAGCGGAGCGGAGCGGAGCGGAGCGGAGCGGAGCGGAGCGGAGCGGAGAGCGGAGAGCGGAGAGCGGAGAGCGGAGAGCGGAGAGCGGAGAGCGGAGAGCGGAGAGCGGAGAGCGGAGAGCGAGGAG$	225
QУ	297	GlyAspLeuAlaGluArgValGluValValAlaLeuSerAspCysGlnGlnGluLeuSer	316
Db	2260	$\tt GGTGACCTGGCAGAAAGGGTCGAGGTGGTCGCCCTGAGTGACTGTCAGCAGGAGTTGAGC$	231
QУ	317	LeuValArgThrValThrArgGlyProArgAlaPheLeuSerArgGluGluAlaGlnHis	336
Db	2320	TTGGTGAGGACTGTGACTCGGGGACCACGCGCCTTCCTGAGCCGCGAGGAGGCACACCAC	237
QУ	337	PheIleLysAlaCysGlyLeuLeuAsnCysGluAlaValLeuGlnLeuLeuThrCysHis	356
Db	2380	$\tt TTCATCAAAGCGTGTGGACTGCTCAACTGTGAGGCCGTGCTGCAGCTGCTGACCTGCCAC$	243
Qу	357	LeuArgGlyThrSerGluCysThrGlnLeuArgAlaLeuCysAlaIleAlaSerLeuGly	376
Db	2440	$\tt CTGCGTGGGACCAGTGAATGCACGCAGCTGAGGGCGCTGTGTGCCATCGCCTCCCTGGGG$	249
QУ	377	$Ser Ser {\tt AspLeuLeuProGlnGluHisIleLeuLeuArgThrArgProTrpLeuGlnGlu}$	396

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Db	2500	${\tt AGCAGCGACCTCCTCCCCCAGGAGCACATCCTCCTCCGCACCCGGCCGTGGCTGCAGGAG}$	255
Qy	397	LeuSerMetGlySerProGlyProValThrAsnLysAlaThrLysIleLeuArgHisPhe	416
Db	2560	CTCAGCATGGGCAGCCCGGGACCTGTGACCAACAAGGCCACCAAGATCCTGAGGCACTTT	261
QУ	417	GluAlaSerCysGlyGlnLeuSerProAlaArgGlyThrSerAlaGluProGlyProThr	436
Db	2620	GAGGCCTCCTGTGGGCAGCTGTCCCCTGCCCGGGGCACCTCAGCCTGAGCCTGGCCCCACA	267
Qу	437	AlaAlaLeuProGlyProSerAspLeuLeuThrAspAlaValProLeuProGlySerGln	456
Db	2680	GCCGCCTCCCAGGCCATCTGACCTGACCGACGCTGTGCCTCTCCCTGGGAGCCAG	273
QУ	457	$Val {\tt PheLeuGlnProLeuSerSerThrProValSerSerArgSerProAlaProSerSer}$	476
Db	2740	GTCTTCCTGCAGCCTCTGAGTTCAACCCCGGTCTCGTCCCGGAGCCCTGCTCCCTCATCT	279
Qу	477	${\tt GlyMetProSerSerProValProThrProProProAspAlaSerProIleProAlaPro}$	496
Db	2800	GGGATGCCGTCCAGCCCTGTGCCCACCCCACCCCAGATGCCTCCCCCATTCCAGCCCCC	285
Qу	497	GlyAspProSerGluAlaGluAlaArgLeuAlaGluSerArgArgTrpArgProGluArg	516
Db	2860	GGAGACCCCAGCGAGGCCAGACTGGCAGAAAGCAGGCGGTGGAGACCTGAACGG	291
QУ	517	${\tt IleProGlyGlyThrAspSerProLysArgGlyProSerSerCysAlaTrpSerArgAsp}$	536
Db	2920	ATCCCGGGGGGCACGGACAGCCCAAAGAGAGCCCCAGCAGCTGTGCGTGGAGCCGCGAC	297
Qу	537	${\tt SerLeuPheAlaGlyMetGluLeuValAlaCysProArgLeuValGlyAlaGlyAlaAla}$	556
Db	2980	TCCTTGTTTGCTGGCATGGAGCTGGTGGCCTGTCCCCGCCTGGTGGGGGCTGGGGCTGCT	303
Qу	557	${\tt AlaGlyGluSerCysProAspAlaProArgAlaProGlnThrSerSerGlnArgThrAla}$	576
Db	3040		309
QУ	577	AlaLysGluProProGlySerGluProSerAlaPheAlaPheLeuAsnAla 593	
Db	3100	GCCAAAGACCTCCTGGCTCAGAGCCGTCAGCTTTCGCGTTCCTGAACGCC 3150	

3. Sequence alignment for AX714043:

```
US-10-537-767-4 (1-593) x AX714043 (1-2249)
        {\tt 1~MetAlaAlaProProLeuArgAspArgLeuSerPheLeuHisArgLeuProIleLeu~20}\\
Qу
          23 ATGGCTGCCGCCGCCGCTACGGGACCGCCTGAGCTTTCTACACCGGCTCCCGATTCTC 82
Db
ΩУ
        {\tt 21\ LeuLysGlyThrSerAspAspValProCysProGlyTyrLeuPheGluGluIleAla\ 40}
          Db
        83 CTGAAGGGGACGTCCGATGATGATGTCCCGTGTCCGGGCTACCTGTTTGAAGAGATTGCT 142
        {\tt 41~LysIleSerHisGluSerProGlySerSerGlnCysLeuLeuGluTyrLeuLeuSerArg~60}
Qу
          143 AAAATCTCCCACGAGTCTCCGGGCAGCAGCCAGTGCCTGCTGGAGTACCTCCTGAGCCGC 202
        61 LeuHisSerSerSerGlyHisGlyLysLeuLysValLeuLysIleLeuLeuTyrLeuCys 80
Qу
          203 CTGCACAGCAGCTCCGGCCACGGGAAGCTCAAGGTGCTGAAGATCCTGCTCTATCTGTGC 262
Db
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QУ	81	SerHisGlySerSerPhePheLeuLeuIleLeuLysArgAsnSerAlaPheIleGlnGlu	100
Db	263	AGCCACGGCTCCTTCTTCCTGCTCATCCTCAAACGCAACTCTGCCTTCATCCAGGAA	322
Qу	101	AlaAlaPheAlaGlyProProAspProLeuHisGlyAsnSerLeuTyrGlnLysVal	120
Db	323	$\tt GCTGCAGCTTTTGCAGGGCCCCCAGATCCTCTGCACGGGAACAGCTTGTACCAGAAGGTT$	382
QУ	121	ArgAlaAlaGlnAspLeuGlySerThrLeuPheSerAspThrValLeuProLeuAla	140
Db	383	$\tt CGCGCGGCGCAGGACTTGGGGAGCACCCTGTTCTCGGACACCGTGTTGCCGCTGGCT$	442
Qу	141	ProSerGlnProLeuGlyThrProProAlaThrGlyMetGlySerGlnAlaArgProHis	160
Db	443		502
QУ	161	SerThrLeuGlnGlyPheGlyTyrSerLysGluHisGlyArgThrGlySerAlaGlyGlu	180
Db	503	AGCACCTCCAGGGTTTCGGCTACAGCAAGGAACACGGCCGCACG	547
Qу	181	$\verb AlaPheLeuSerThrIleGlnLysAlaAlaGluValValAlaSerAlaMetArgProGly $	200
Db	547		547
Qу	201	${\tt ProGluSerProSerThrArgArgLeuLeuProArgGlyAspThrTyrGlnProAlaMet}$	220
Db	547		547
QУ	221	${\tt MetProSerAlaSerHisGlyProProThrLeuGlyAsnLeuLeuProGlyAlaIlePro}$	240
Db	547		547
Qу	241	${\tt GlyProArgAlaValArgHisGlnProGlyGlnAlaGlyGlyGlyTrpAspGluLeuAsp}$	260
Db	548	GCTGTGAGGCATCAGCCTGGGCAGGCCGGAGGGGGCTGGGATGAGCTGGAC	598
Qу	261	Ser Gly ProSer Ser Gln Asn Ser Ser Gln Asn Ser Asp Leu Ser Arg Val Ser Asp Ser Asp Leu Ser Asp Leu Ser Asp Leu Ser Asp Ser Asp Leu Ser Asp Leu Ser Asp Ser Asp Leu Ser Asp Leu Ser Asp Leu Ser Asp Ser Asp Leu Ser A	280
Db	599	AGCGGCCCCAGCTCTCAGAATTCCTCCCAGAACAGTGACCTGAGCAGGGTCTCGGACTCG	658
Qу	281	${\tt GlySerHisSerGlySerAspSerHisSerGlyAlaSerArgGluProGlyAspLeuAla}$	300
Db	659	GGCAGTCATTCCGGCAGCGACAGCCATTCAGGGGCCAGCCGGGAGCCGGGTGACCTGGCA	718
Qу	301	${\tt GluArgValGluValValAlaLeuSerAspCysGlnGlnGluLeuSerLeuValArgThr}$	320
Db	719	GAAAGGGTCGAGTGGCCCTGAGTGACTGTCAGCAGGAGTTGAGCTTGGTGAGGACT	778
Qу	321	ValThrArgGlyProArgAlaPheLeuSerArgGluGluAlaGlnHisPheIleLysAla	
Db	779	GTGACTCGGGGACCACGCGCTTCCTGAGCCGCGAGGAGGCACAGCACTTCATCAAAGCG	
Qу	341	${\tt CysGlyLeuLeuAsnCysGluAlaValLeuGlnLeuLeuThrCysHisLeuArgGlyThr}$	360
Db	839	TGTGGACTGCTCAACTGTGAGGCCGTGCTGCAGCTGCTGACCTGCCACCTGCGTGGGACC	898
Qу	361	Ser Glu Cys Thr Gln Leu Arg Ala Leu Cys Ala Ile Ala Ser Leu Gly Ser Ser Asp Leu Cys Ala Ile Ala Ser Leu Gly Ser Ser Asp Leu Cys Ala Ile Ala Ser Leu Gly Ser Asp Leu Cys Asp	380
Db	899	AGTGAATGCACGCAGCTGAGGGCGCTGTGTGCCATCGCCTCCCTGGGGAGCAGCGACCTC	958
QУ	381	Leu Pro Gln Glu His Ile Leu Leu Arg Thr Arg Pro Trp Leu Gln Glu Leu Ser Met Gly Glu Leu Gln Glu	400
Db	959		101
Qу	401	SerProGlyProValThrAsnLysAlaThrLysIleLeuArgHisPheGluAlaSerCys	420
Dh	1010	ACCCCCCCACCTACCTACCAACAACAACCAACCAACATCCTCACCCACTTTTCACCCCTCCT	107

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421 GlyGlnLeuSerProAlaArgGlyThrSerAlaGluProGlyProThrAlaAlaLeuPro 440
QУ
          1079 GGGCAGCTGTCCCCTGCCGGGGCACCTCAGCTGAGCCTGGCCCCACAGCCGCCCTCCCA 1138
Dh
       441 GlyProSerAspLeuLeuThrAspAlaValProLeuProGlySerGlnValPheLeuGln 460
Qv
          1139 GGCCCATCTGACCTGCTGACCGACGCTGTGCCTCCCTGGGAGCCAGGTCTTCCTGCAG 1198
0v
       461 ProLeuSerSerThrProValSerSerArgSerProAlaProSerSerGlyMetProSer 480
          Db
      1199 CCTCTGAGTTCAACCCCGGTCTCGTCCCGGAGCCCTGCTCCCTCATCTGGGATGCCGTCC 1258
       481 SerProValProThrProProProAspAlaSerProIleProAlaProGlyAspProSer 500
          1259 AGCCCTGTGCCCACCCCACACTCCAGATGCCTCCCCATTCCAGCCCCCGGAGACCCCAGC 1318
Db
       501 GluAlaGluAlaArgLeuAlaGluSerArgArgTrpArgProGluArgIleProGlyGly 520
Qy
          1319 GAGGCCGAGGCCAGACTGGCAGAAAGCAGGCGGTGGAGACCTGAACGGATCCCAGGGGGC 1378
Db
       521 ThrAspSerProLysArgGlyProSerSerCysAlaTrpSerArgAspSerLeuPheAla 540
0.7
          1379 ACGGACAGCCCAAAGAGAGCCCCAGCAGCTGTGCGTGGAGCCGCGACTCCTTGTTTGCT 1438
       541 GlyMetGluLeuValAlaCysProArgLeuValGlyAlaGlyAlaAlaAlaGlyGluSer 560
QУ
          1439 GGCATGGAGCTGGTGGCCTGTCCCCGCCTGGTGGGGGCTGCTGCGGGGAGAGTCC 1498
       561 CysProAspAlaProArgAlaProGlnThrSerSerGlnArgThrAlaAlaLysGluPro 580
Qy
          1499 TGTCCTGATGCTCCCCGCGCCCCCAAACATCGTCCCAGAGGACAGCAGCCAAAGAGCCT 1558
Db
       581 ProGlySerGluProSerAlaPheAlaPheLeuAsnAla 593
QУ
          1559 CCTGGCTCAGAGCCGTCAGCTTTCGCGTTCCTGAACGCC 1597
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Claims 1-5 are allowed.

Any comments considered necessary by applicant must be submitted no later than the payment of the issue fee and, to avoid processing delays, should preferably accompany the issue fee. Such submissions should be clearly labeled "Comments on Statement of Reasons for Allowance."

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Jae W. Lee whose telephone number is 571-272-9949. The examiner can normally be reached between 9:00 to 5:00 on Monday-Friday.

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If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Kathleen Kerr Bragdon can be reached on 571-272-0931. The fax phone number for the organization where this application or proceeding is assigned is 571-

273-8300.

Information regarding the status of an application may be obtained from the Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free). If you would like assistance from a USPTO Customer Service Representative or access to the automated information system, call

/JAE W LEE/

Examiner, Art Unit 1656

/Delia M. Ramirez/

Delia M. Ramirez, Ph.D.

Primary Examiner, Art Unit 1652

800-786-9199 (IN USA OR CANADA) or 571-272-1000.